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methylation ("whole genome amplification") (c) [Advanced Scholar Search](#)

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In vitro DNA immortalization and whole genome amplification using libraries generated from randomly fragmented DNA

JH Pinter, T Kurihara, I Sieptsova ... - US Patent App. 10/ ... 2004 - Google Patents
... (43) Pub. Date: Oct. 21, 2004 (54) IN VITRO DNA IMMORTALIZATION AND **WHOLE GENOME AMPLIFICATION** USING LIBRARIES GENERATED FROM RANDOMLY FRAGMENTED DNA
Related US Application Data (60) Provisional application No. 60/453,071, filed on Mar. ...
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The Hematopoietic System-specific Minor Histocompatibility Antigen HA-1 Shows Aberrant Expression in Epithelial Cancer Cells

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CA Klein, M Wilke, J Pool, C Vermeulen ... - The Journal of ... 2002 - jern.rupress.org
... HA-1, CD45, and PBGD expression were quantified in all test samples using these **calibration** functions. ... The isolated DNA was subjected to **whole genome amplification** and CGH. ... DNA **methylation** has been proposed as the primary mechanism for the inactivation of genes (24). ...
[Cited by 75](#) - [Related articles](#) - [Bib. Direct](#) - [All 11 versions](#)

Multiplexed Methylation Detection Methods

JB Fan - US Patent App. 10/537,204, 2003 - Google Patents
... Date: Nov. 22, 2007 (54) MULTIPLEXED **METHYLATION** DETECTION METHODS (76) Inventor: Jian-Bing Fan, San Diego, CA (US) Correspondence Address: MCDERMOTT, WILL & EMERY
4370 LA JOLLA VILLAGE DRIVE, SUITE 700 SAN DIEGO, CA 92122 (US) (21) Appl. ...

Multiplexed Methylation Detection Methods

J Fan - US Patent App. 20,070,269, 2003 - freepatentsonline.com
... The invention also provides a **calibration** curve comprising a first axis indicating the amount of unmethylated template population compared to methylated template ... The invention methods for detecting **methylation** allow for **whole genome amplification** after bisulfite ...
[Cached](#)

The San Diego Conference Nucleic Acid Technology: The Cutting Edge of Discovery November 6-8, 1997

[\[HTML\] from clinchem.org](#)

S Graham - Clinical Chemistry 1997 - Am Assoc Clin Chem
... To address these limitations, we have combined **whole genome amplification** by primer extension ... the "competitive" nature of the amplification of the target and quantitation **standard**. ... Regulatory region mutations as well as **methylation** represent additional possible mechanisms ...

Polynucleotide sequence detection assays

BB Rosenblum - US Patent App. 20,050,053, 2003 - freepatentsonline.com
... addition, preparation of target nucleic acids can be accomplished using **whole genome amplification** techniques (eg ... also be used to estimate the average amount of **methylation** for one ... by including nucleotide analogs having higher binding affinities than **standard** nucleotides for ...
[Cached](#)

Polynucleotide sequence detection assays

BB Rosenblum - US Patent App. 10/718,016, 2003 - Google Patents
... of target nucleic acids can be accomplished using **whole genome amplification** techniques (eg, Lizardi, US Pat. No. 6,124,120). [0041] In some embodiments, target nucleic acids are chemically treated prior to analysis. For example, analysis of the **methylation** state of cytosines ...

Behavioral and metabolic consequences of neonatal exposure to diazepam in rat pups

H Schroeder, AC Humbert, V Kotzki, D Desor ... - Experimental ... 1995 - Elsevier
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Cellular engineering, protein expression profiling, differential labeling of peptides, and novel reagents therefor

JM Short, M Lafferich, J Wei ... - US Patent App. 20,030/ ... 2002 - freepatentsonline.com
... labeling reagents, which relied on stable isotopes, which are expensive, and not flexible to differential labeling of more than two mixtures of peptides; labeling methods limited only to **methylation** of carboxy ... **Standard** IUPAC nomenclature is used for describing pooled nucleotides ...
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methylation ("whole genome amplification")

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